

SCORE Search Results Details for Application 10552515 and Search Result 20090316_112516_us-10-552-515-3.ra1.

| | | | | |
|----------------------------|--------------------------------------|------------------------------|-----------------------|-----------------------------|
| Score Home | Retrieve Application | SCORE System | SCORE | Comments / |
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This page gives you Search Results detail for the Application 10552515 and Search Result 20090316_112516_us-10-552-515-3.ra1.

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GenCore version 6.3

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OM protein - protein search, using sw model

Run on: March 17, 2009, 05:01:40 ; Search time 2 Seconds
(without alignments)
1258.128 Million cell updates/sec

Title: US-10-552-515-3
Perfect score: 46
Sequence: 1 SLFMALWAV 9

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 1316349 seqs, 215321474 residues

Total number of hits satisfying chosen parameters: 1316349

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Issued_Patents_AA:*
1: /ABSS/Data/CRF/ptodata/1/iaa/5_COMB.pep:*
2: /ABSS/Data/CRF/ptodata/1/iaa/6_COMB.pep:*
3: /ABSS/Data/CRF/ptodata/1/iaa/7_COMB.pep:*
4: /ABSS/Data/CRF/ptodata/1/iaa/H_COMB.pep:*
5: /ABSS/Data/CRF/ptodata/1/iaa/PCTUS_COMB.pep:*
6: /ABSS/Data/CRF/ptodata/1/iaa/RE_COMB.pep:*
7: /ABSS/Data/CRF/ptodata/1/iaa/backfiles1.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

%

| Result No. | Score | Query Match | Length | DB | ID | Description |
|------------|-------|-------------|--------|----|----------------------|-------------------|
| 1 | 40 | 87.0 | 117 | 3 | US-10-703-032-142336 | Sequence 142336, |
| 2 | 39 | 84.8 | 642 | 3 | US-10-108-260A-4483 | Sequence 4483, Ap |
| 3 | 39 | 84.8 | 956 | 3 | US-10-912-745B-284 | Sequence 284, App |
| 4 | 36 | 78.3 | 207 | 2 | US-08-811-519-30 | Sequence 30, Appl |
| 5 | 36 | 78.3 | 220 | 2 | US-09-489-039A-13425 | Sequence 13425, A |
| 6 | 36 | 78.3 | 250 | 2 | US-09-248-796A-20183 | Sequence 20183, A |
| 7 | 36 | 78.3 | 274 | 4 | US-10-038-895A-1 | Sequence 1, Appli |
| 8 | 36 | 78.3 | 440 | 2 | US-09-631-603-22 | Sequence 22, Appl |
| 9 | 36 | 78.3 | 440 | 2 | US-09-826-509-567 | Sequence 567, App |
| 10 | 36 | 78.3 | 440 | 3 | US-10-925-095-567 | Sequence 567, App |
| 11 | 36 | 78.3 | 440 | 3 | US-11-404-939-567 | Sequence 567, App |
| 12 | 36 | 78.3 | 442 | 2 | US-09-538-092-637 | Sequence 637, App |
| 13 | 36 | 78.3 | 449 | 1 | US-08-142-439A-5 | Sequence 5, Appli |
| 14 | 36 | 78.3 | 449 | 1 | US-08-869-477-5 | Sequence 5, Appli |
| 15 | 35 | 76.1 | 487 | 2 | US-09-328-352-6206 | Sequence 6206, Ap |
| 16 | 34 | 73.9 | 38 | 3 | US-10-105-299-3648 | Sequence 3648, Ap |
| 17 | 34 | 73.9 | 108 | 2 | US-09-489-039A-13025 | Sequence 13025, A |
| 18 | 34 | 73.9 | 144 | 3 | US-10-703-032-126625 | Sequence 126625, |
| 19 | 34 | 73.9 | 152 | 2 | US-09-489-039A-11538 | Sequence 11538, A |
| 20 | 34 | 73.9 | 218 | 2 | US-09-270-767-42075 | Sequence 42075, A |
| 21 | 34 | 73.9 | 435 | 2 | US-09-252-991A-19124 | Sequence 19124, A |
| 22 | 34 | 73.9 | 574 | 3 | US-10-912-745B-229 | Sequence 229, App |
| 23 | 34 | 73.9 | 574 | 3 | US-10-912-745B-230 | Sequence 230, App |
| 24 | 34 | 73.9 | 968 | 3 | US-09-252-691C-7784 | Sequence 7784, Ap |
| 25 | 33 | 71.7 | 15 | 3 | US-11-129-741A-1158 | Sequence 1158, Ap |
| 26 | 33 | 71.7 | 169 | 2 | US-10-094-749-1824 | Sequence 1824, Ap |
| 27 | 33 | 71.7 | 202 | 3 | US-10-703-032-125681 | Sequence 125681, |
| 28 | 33 | 71.7 | 225 | 3 | US-09-540-209B-7498 | Sequence 7498, Ap |
| 29 | 33 | 71.7 | 240 | 3 | US-10-703-032-136346 | Sequence 136346, |
| 30 | 33 | 71.7 | 252 | 3 | US-10-369-493-7925 | Sequence 7925, Ap |
| 31 | 33 | 71.7 | 362 | 3 | US-10-369-493-4227 | Sequence 4227, Ap |
| 32 | 33 | 71.7 | 469 | 2 | US-09-328-352-5007 | Sequence 5007, Ap |
| 33 | 33 | 71.7 | 507 | 3 | US-10-369-493-10701 | Sequence 10701, A |
| 34 | 33 | 71.7 | 524 | 2 | US-09-252-991A-18580 | Sequence 18580, A |
| 35 | 33 | 71.7 | 528 | 3 | US-09-602-740-34 | Sequence 34, Appl |
| 36 | 33 | 71.7 | 528 | 3 | US-10-781-014-34 | Sequence 34, Appl |
| 37 | 33 | 71.7 | 530 | 3 | US-09-602-740-32 | Sequence 32, Appl |
| 38 | 33 | 71.7 | 530 | 3 | US-10-781-014-32 | Sequence 32, Appl |
| 39 | 33 | 71.7 | 575 | 3 | US-10-805-394A-4263 | Sequence 4263, Ap |
| 40 | 33 | 71.7 | 595 | 3 | US-10-703-032-120251 | Sequence 120251, |
| 41 | 33 | 71.7 | 596 | 2 | US-10-104-047-2541 | Sequence 2541, Ap |
| 42 | 33 | 71.7 | 920 | 2 | US-10-104-047-2574 | Sequence 2574, Ap |
| 43 | 33 | 71.7 | 1280 | 3 | US-10-343-657-7 | Sequence 7, Appli |
| 44 | 33 | 71.7 | 1359 | 3 | US-10-736-769-44 | Sequence 44, Appl |
| 45 | 32 | 69.6 | 86 | 3 | US-10-198-232-64 | Sequence 64, Appl |

ALIGNMENTS

RESULT 1

US-10-703-032-142336

; Sequence 142336, Application US/10703032

```

; Patent No. 7214786
; GENERAL INFORMATION:
; APPLICANT: Kovalic, David K.
; APPLICANT: Andersen, Scott E.
; APPLICANT: Byrum, Joseph R.
; APPLICANT: Conner, Timothy W.
; APPLICANT: Cao, Yongwei
; APPLICANT: Masucci, James D.
; APPLICANT: Zhou, Yihua
; TITLE OF INVENTION: Nucleic Acid Molecules And Other Molecules Associated With
; TITLE OF INVENTION: Plants
; FILE REFERENCE: 38-21(53374)B
; CURRENT APPLICATION NUMBER: US/10/703,032
; CURRENT FILING DATE: 2003-11-06
; PRIOR APPLICATION NUMBER: 10/020,338
; PRIOR FILING DATE: 2001-12-12
; NUMBER OF SEQ ID NOS: 211164
; SEQ ID NO 142336
; LENGTH: 117
; TYPE: PRT
; ORGANISM: Triticum aestivum
; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_TA_36754.pep
US-10-703-032-142336

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Query Match          87.0%; Score 40; DB 3; Length 117;
Best Local Similarity 77.8%; Pred. No. 21;
Matches      7; Conservative      2; Mismatches      0; Indels      0; Gaps      0;

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Qy      1 SLFMALWAV 9
        |:|:|||||
Db      96 SIFIALWAV 104

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RESULT 2

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US-10-108-260A-4483
; Sequence 4483, Application US/10108260A
; Patent No. 7193069
; GENERAL INFORMATION:
; APPLICANT: HELIX RESEARCH INSTITUTE
; TITLE OF INVENTION: No. 7193069e1 full length cDNA
; FILE REFERENCE: H1-A0106
; CURRENT APPLICATION NUMBER: US/10/108,260A
; CURRENT FILING DATE: 2002-03-27
; NUMBER OF SEQ ID NOS: 5458
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 4483
; LENGTH: 642
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-108-260A-4483

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Query Match          84.8%; Score 39; DB 3; Length 642;
Best Local Similarity 87.5%; Pred. No. 1.7e+02;
Matches      7; Conservative      1; Mismatches      0; Indels      0; Gaps      0;

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Qy 1 SLFMALWA 8
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 Db 385 SVFMALWA 392

RESULT 3

US-10-912-745B-284
 ; Sequence 284, Application US/10912745B
 ; Patent No. 7473531
 ; GENERAL INFORMATION
 ; APPLICANT: DOMON, Bruno et al.
 ; TITLE OF INVENTION: Pancreatic Cancer Targets and Uses
 ; TITLE OF INVENTION: Thereof
 ; FILE REFERENCE: CL001538
 ; CURRENT APPLICATION NUMBER: US/10/912,745B
 ; CURRENT FILING DATE: 2004-08-06
 ; NUMBER OF SEQ ID NOS: 875
 ; SOFTWARE: FastSEQ for Windows Version 4.0
 ; SEQ ID NO 284
 ; LENGTH: 956
 ; TYPE: PRT
 ; ORGANISM: Homo sapiens
 US-10-912-745B-284

Query Match 84.8%; Score 39; DB 3; Length 956;
 Best Local Similarity 87.5%; Pred. No. 2.5e+02;
 Matches 7; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1 SLFMALWA 8
 |:|||||
 Db 413 SVFMALWA 420

RESULT 4

US-08-811-519-30
 ; Sequence 30, Application US/08811519B
 ; Patent No. 6630345
 ; GENERAL INFORMATION:
 ; APPLICANT: Petrenko, Alexandre
 ; TITLE OF INVENTION: CALCIUM INDEPENDENT RECEPTOR OF ALPHA-LATROTOXIN,
 ; TITLE OF INVENTION: CHARACTERIZATION AND USES THEREOF
 ; FILE REFERENCE: 1049-1-007
 ; CURRENT APPLICATION NUMBER: US/08/811,519B
 ; CURRENT FILING DATE: 1997-03-04
 ; NUMBER OF SEQ ID NOS: 31
 ; SOFTWARE: PatentIn Ver. 2.0
 ; SEQ ID NO 30
 ; LENGTH: 207
 ; TYPE: PRT
 ; ORGANISM: rat
 US-08-811-519-30

Query Match 78.3%; Score 36; DB 2; Length 207;
 Best Local Similarity 55.6%; Pred. No. 1.8e+02;
 Matches 5; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

Qy 1 SLFMALWAV 9
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 Db 110 AIFVALWAI 118

RESULT 5

US-09-489-039A-13425
 ; Sequence 13425, Application US/09489039A
 ; Patent No. 6610836
 ; GENERAL INFORMATION:
 ; APPLICANT: Gary Breton et. al
 ; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA
 ; TITLE OF INVENTION: PNEUMONIAE FOR DIAGNOSTICS AND THERAPEUTICS
 ; FILE REFERENCE: 2709.2004001
 ; CURRENT APPLICATION NUMBER: US/09/489,039A
 ; CURRENT FILING DATE: 2000-01-27
 ; PRIOR APPLICATION NUMBER: US 60/117,747
 ; PRIOR FILING DATE: 1999-01-29
 ; NUMBER OF SEQ ID NOS: 14342
 ; SEQ ID NO 13425
 ; LENGTH: 220
 ; TYPE: PRT
 ; ORGANISM: Klebsiella pneumoniae
 US-09-489-039A-13425

Query Match 78.3%; Score 36; DB 2; Length 220;
 Best Local Similarity 77.8%; Pred. No. 1.9e+02;
 Matches 7; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 1 SLFMALWAV 9
 |||| || |
 Db 9 SLFMKLWLV 17

RESULT 6

US-09-248-796A-20183
 ; Sequence 20183, Application US/09248796A
 ; Patent No. 6747137
 ; GENERAL INFORMATION:
 ; APPLICANT: Keith Weinstock et al
 ; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICANS
 ; TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
 ; FILE REFERENCE: 107196.132
 ; CURRENT APPLICATION NUMBER: US/09/248,796A
 ; CURRENT FILING DATE: 1999-02-12
 ; PRIOR APPLICATION NUMBER: US 60/074,725
 ; PRIOR FILING DATE: 1998-02-13
 ; PRIOR APPLICATION NUMBER: US 60/096,409
 ; PRIOR FILING DATE: 1998-08-13
 ; NUMBER OF SEQ ID NOS: 28208
 ; SEQ ID NO 20183
 ; LENGTH: 250
 ; TYPE: PRT
 ; ORGANISM: Candida albicans

US-09-248-796A-20183

Query Match 78.3%; Score 36; DB 2; Length 250;
 Best Local Similarity 77.8%; Pred. No. 2.1e+02;
 Matches 7; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy 1 SLFMALWAV 9
 ||:|||||
 Db 70 SLIIALWAV 78

RESULT 7

US-10-038-895A-1

; Sequence 1, Application US/10038895A
 ; Patent No. H002136
 ; GENERAL INFORMATION:
 ; APPLICANT: Kulp, David C.
 ; APPLICANT: Siani-Rose, Michael A.
 ; APPLICANT: Williams, Alan J.
 ; APPLICANT: Harmon, Cyrus L.
 ; TITLE OF INVENTION: Nucleic Acids Encoding G Proteins Coupled Receptors
 ; FILE REFERENCE: 3379.1
 ; CURRENT APPLICATION NUMBER: US/10/038,895A
 ; CURRENT FILING DATE: 2003-03-25
 ; PRIOR APPLICATION NUMBER: US 60/244,082
 ; PRIOR FILING DATE: 2000-10-26
 ; NUMBER OF SEQ ID NOS: 20
 ; SOFTWARE: PatentIn version 3.2
 ; SEQ ID NO 1
 ; LENGTH: 274
 ; TYPE: PRT
 ; ORGANISM: Artificial Sequence
 ; FEATURE:
 ; OTHER INFORMATION: Synthetic Organism
 ; FEATURE:
 ; NAME/KEY: misc_feature
 ; LOCATION: (126)..(126)
 ; OTHER INFORMATION: Xaa can be any naturally occurring amino acid
 ; FEATURE:
 ; NAME/KEY: misc_feature
 ; LOCATION: (146)..(146)
 ; OTHER INFORMATION: Xaa can be any naturally occurring amino acid
 US-10-038-895A-1

Query Match 78.3%; Score 36; DB 4; Length 274;
 Best Local Similarity 55.6%; Pred. No. 2.3e+02;
 Matches 5; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

Qy 1 SLFMALWAV 9
 ::|:||||:
 Db 170 AIFVALWAI 178

RESULT 8

US-09-631-603-22

; Sequence 22, Application US/09631603

```

; Patent No. 6733990
; GENERAL INFORMATION:
; APPLICANT: Hodge, Martin R.
; APPLICANT: Lloyd, Clare
; APPLICANT: Weich, Nadine
; TITLE OF INVENTION: 15571, A No. 6733990el GPCR-like Molecule of the
; TITLE OF INVENTION: Secretin-Like Family and Uses Thereof
; FILE REFERENCE: 5800-48A
; CURRENT APPLICATION NUMBER: US/09/631,603
; CURRENT FILING DATE: 2000-08-03
; PRIOR APPLICATION NUMBER: 09/515,781
; PRIOR FILING DATE: 2000-02-29
; PRIOR APPLICATION NUMBER: 60/146,916
; PRIOR FILING DATE: 2000-08-03
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 22
; LENGTH: 440
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-631-603-22

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Query Match      78.3%; Score 36; DB 2; Length 440;
Best Local Similarity 55.6%; Pred. No. 3.7e+02;
Matches      5; Conservative      4; Mismatches      0; Indels      0; Gaps      0;

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Qy      1 SLFMALWAV 9
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Db      267 AIFVALWAI 275

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RESULT 9
US-09-826-509-567
; Sequence 567, Application US/09826509
; Patent No. 6806054
; GENERAL INFORMATION:
; APPLICANT: Lehmann-Bruinsma, Karin
; APPLICANT: Liaw, Chen W.
; APPLICANT: Lin, I-Lin
; TITLE OF INVENTION: No. 6806054-Endogenous, Constitutively Activated Known G
; TITLE OF INVENTION: Protein-Coupled Receptors
; FILE REFERENCE: AREN-207
; CURRENT APPLICATION NUMBER: US/09/826,509
; CURRENT FILING DATE: 2001-04-05
; PRIOR APPLICATION NUMBER: 60/195,747
; PRIOR FILING DATE: 2000-04-07
; PRIOR APPLICATION NUMBER: 09/170,496
; PRIOR FILING DATE: 1998-10-13
; NUMBER OF SEQ ID NOS: 589
; SOFTWARE: PatentIn Version 2.1
; SEQ ID NO 567
; LENGTH: 440
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-826-509-567

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Query Match 78.3%; Score 36; DB 2; Length 440;
 Best Local Similarity 55.6%; Pred. No. 3.7e+02;
 Matches 5; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

Qy 1 SLFMALWAV 9
 ::|:||||:
 Db 267 AIFVALWAI 275

RESULT 10

US-10-925-095-567
 ; Sequence 567, Application US/10925095
 ; Patent No. 7097969
 ; GENERAL INFORMATION:
 ; APPLICANT: Lehmann-Bruinsma, Karin
 ; APPLICANT: Liaw, Chen W.
 ; APPLICANT: Lin, I-Lin
 ; TITLE OF INVENTION: No. 7097969-Endogenous, Constitutively Activated Known G
 ; TITLE OF INVENTION: Protein-Coupled Receptors
 ; FILE REFERENCE: AREN-207
 ; CURRENT APPLICATION NUMBER: US/10/925,095
 ; CURRENT FILING DATE: 2004-08-24
 ; PRIOR APPLICATION NUMBER: US/09/826,509
 ; PRIOR FILING DATE: 2001-04-05
 ; PRIOR APPLICATION NUMBER: 60/195,747
 ; PRIOR FILING DATE: 2000-04-07
 ; PRIOR APPLICATION NUMBER: 09/170,496
 ; PRIOR FILING DATE: 1998-10-13
 ; NUMBER OF SEQ ID NOS: 589
 ; SOFTWARE: PatentIn Version 2.1
 ; SEQ ID NO 567
 ; LENGTH: 440
 ; TYPE: PRT
 ; ORGANISM: Homo sapiens
 US-10-925-095-567

Query Match 78.3%; Score 36; DB 3; Length 440;
 Best Local Similarity 55.6%; Pred. No. 3.7e+02;
 Matches 5; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

Qy 1 SLFMALWAV 9
 ::|:||||:
 Db 267 AIFVALWAI 275

RESULT 11

US-11-404-939-567
 ; Sequence 567, Application US/11404939
 ; Patent No. 7381522
 ; GENERAL INFORMATION:
 ; APPLICANT: Lehmann-Bruinsma, Karin
 ; APPLICANT: Liaw, Chen W.
 ; APPLICANT: Lin, I-Lin
 ; TITLE OF INVENTION: Non-Endogenous, Constitutively Activated Known G
 ; TITLE OF INVENTION: Protein-Coupled Receptors


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; FILE REFERENCE: AREN-207
; CURRENT APPLICATION NUMBER: US/11/404,939
; CURRENT FILING DATE: 2006-04-14
; PRIOR APPLICATION NUMBER: US/09/826,509
; PRIOR FILING DATE: 2001-04-05
; PRIOR APPLICATION NUMBER: 60/195,747
; PRIOR FILING DATE: 2000-04-07
; PRIOR APPLICATION NUMBER: 09/170,496
; PRIOR FILING DATE: 1998-10-13
; NUMBER OF SEQ ID NOS: 589
; SOFTWARE: PatentIn Version 2.1
; SEQ ID NO 567
; LENGTH: 440
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-404-939-567

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Query Match          78.3%;  Score 36;  DB 3;  Length 440;
Best Local Similarity 55.6%;  Pred. No. 3.7e+02;
Matches      5;  Conservative      4;  Mismatches      0;  Indels      0;  Gaps      0;

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Qy      1 SLFMALWAV 9
        ::|:||||:
Db      267 AIFVALWAI 275

```

RESULT 12

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US-09-538-092-637
; Sequence 637, Application US/09538092
; Patent No. 6753314
; GENERAL INFORMATION:
; APPLICANT: Giot, Loic
; APPLICANT: Mansfield, Traci A.
; TITLE OF INVENTION: Protein-Protein Complexes and Method of Using Same
; FILE REFERENCE: 15966-542
; CURRENT APPLICATION NUMBER: US/09/538,092
; CURRENT FILING DATE: 2000-03-29
; PRIOR APPLICATION NUMBER: 60/127,352
; PRIOR FILING DATE: 1999-04-01
; PRIOR APPLICATION NUMBER: 60/178,965
; PRIOR FILING DATE: 2000-02-01
; NUMBER OF SEQ ID NOS: 1387
; SOFTWARE: CuraPatSeqFormatter Version 0.9
; SEQ ID NO 637
; LENGTH: 442
; TYPE: PRT
; ORGANISM: Saccharomyces cerevisiae
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (0)...(0)
; OTHER INFORMATION: Polypeptide Accession Number YMR243C
US-09-538-092-637

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Query Match          78.3%;  Score 36;  DB 2;  Length 442;
Best Local Similarity 77.8%;  Pred. No. 3.7e+02;

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Matches 7; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy 1 SLFMALWAV 9
 || :|||||
 Db 48 SLLVALWAV 56

RESULT 13

US-08-142-439A-5

; Sequence 5, Application US/08142439A

; Patent No. 5670360

; GENERAL INFORMATION:

; APPLICANT: Thorens, Bernard

; TITLE OF INVENTION: Receptor for the Glucagon-Like-Peptide-1

; TITLE OF INVENTION: (GLP-1)

; NUMBER OF SEQUENCES: 9

; CORRESPONDENCE ADDRESS:

; ADDRESSEE: No. 5670360o No. 5670360disk of No. 5670360th America, Inc.

; STREET: 405 Lexington Avenue, Suite 6400

; CITY: New York

; STATE: New York

; COUNTRY: U.S.A.

; ZIP: 10174-6201

; COMPUTER READABLE FORM:

; MEDIUM TYPE: Floppy disk

; COMPUTER: IBM PC compatible

; OPERATING SYSTEM: PC-DOS/MS-DOS

; SOFTWARE: PatentIn Release #1.0, Version #1.25

; CURRENT APPLICATION DATA:

; APPLICATION NUMBER: US/08/142,439A

; FILING DATE: 24-NOV-93

; CLASSIFICATION: 530

; PRIOR APPLICATION DATA:

; APPLICATION NUMBER: DK 398/92

; FILING DATE: 25-MAR-92

; PRIOR APPLICATION DATA:

; APPLICATION NUMBER: PCT/EP93/00697

; FILING DATE: 23-MAR-93

; ATTORNEY/AGENT INFORMATION:

; NAME: Harrington, James J.

; REGISTRATION NUMBER: 38,711

; REFERENCE/DOCKET NUMBER: 3756.204-US

; TELECOMMUNICATION INFORMATION:

; TELEPHONE: 212 867 0123

; TELEFAX: 212 867 0298

; INFORMATION FOR SEQ ID NO: 5:

; SEQUENCE CHARACTERISTICS:

; LENGTH: 449 amino acids

; TYPE: amino acid

; STRANDEDNESS: single

; TOPOLOGY: linear

; MOLECULE TYPE: protein

; HYPOTHETICAL: NO

; ANTI-SENSE: NO

; ORIGINAL SOURCE:

; ORGANISM: Rattus norvegicus

; STRAIN: Sprague-Dawley
US-08-142-439A-5

Query Match 78.3%; Score 36; DB 1; Length 449;
Best Local Similarity 55.6%; Pred. No. 3.8e+02;
Matches 5; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

Qy 1 SLFMALWAV 9
::|:||||:
Db 267 AIFVALWAI 275

RESULT 14

US-08-869-477-5

; Sequence 5, Application US/08869477
; Patent No. 5846747

; GENERAL INFORMATION:

; APPLICANT: Thorens, Bernard
; TITLE OF INVENTION: Receptor for the Glucagon-Like-Peptide-1
; TITLE OF INVENTION: (GLP-1)
; NUMBER OF SEQUENCES: 9

; CORRESPONDENCE ADDRESS:

; ADDRESSEE: No. 5846747o No. 5846747disk of No. 5846747th America, Inc.
; STREET: 405 Lexington Avenue, Suite 6400
; CITY: New York
; STATE: New York
; COUNTRY: U.S.A.
; ZIP: 10174-6201

; COMPUTER READABLE FORM:

; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25

; CURRENT APPLICATION DATA:

; APPLICATION NUMBER: US/08/869,477
; FILING DATE:
; CLASSIFICATION: 435

; PRIOR APPLICATION DATA:

; APPLICATION NUMBER: US/08/142,439
; FILING DATE: 24-NOV-93
; APPLICATION NUMBER: DK 398/92
; FILING DATE: 25-MAR-92

; PRIOR APPLICATION DATA:

; APPLICATION NUMBER: PCT/EP93/00697
; FILING DATE: 23-MAR-93

; ATTORNEY/AGENT INFORMATION:

; NAME: Harrington, James J.
; REGISTRATION NUMBER: 38,711
; REFERENCE/DOCKET NUMBER: 3756.204-US

; TELECOMMUNICATION INFORMATION:

; TELEPHONE: 212 867 0123
; TELEFAX: 212 867 0298

; INFORMATION FOR SEQ ID NO: 5:

; SEQUENCE CHARACTERISTICS:
; LENGTH: 449 amino acids

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;   TYPE: amino acid
;   STRANDEDNESS: single
;   TOPOLOGY: linear
;   MOLECULE TYPE: protein
;   HYPOTHETICAL: NO
;   ANTI-SENSE: NO
;   ORIGINAL SOURCE:
;   ORGANISM: Rattus norvegicus
;   STRAIN: Sprague-Dawley
US-08-869-477-5

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Query Match          78.3%; Score 36; DB 1; Length 449;
Best Local Similarity 55.6%; Pred. No. 3.8e+02;
Matches      5; Conservative      4; Mismatches      0; Indels      0; Gaps      0;

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Qy      1 SLFMALWAV 9
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Db      267 AIFVALWAI 275

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RESULT 15

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US-09-328-352-6206
; Sequence 6206, Application US/09328352
; Patent No. 6562958
; GENERAL INFORMATION:
; APPLICANT: Gary L. Breton et al.
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ACINETOBACTER
; TITLE OF INVENTION: BAUMANNII FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: GTC99-03PA
; CURRENT APPLICATION NUMBER: US/09/328,352
; CURRENT FILING DATE: 1999-06-04
; NUMBER OF SEQ ID NOS: 8252
; SEQ ID NO 6206
; LENGTH: 487
; TYPE: PRT
; ORGANISM: Acinetobacter baumannii
US-09-328-352-6206

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Query Match          76.1%; Score 35; DB 2; Length 487;
Best Local Similarity 85.7%; Pred. No. 6e+02;
Matches      6; Conservative      1; Mismatches      0; Indels      0; Gaps      0;

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Qy      1 SLFMALW 7
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Db      54 SLFMSLW 60

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Search completed: March 17, 2009, 05:04:35
Job time : 1.76252 secs

SCORE 3.0